

# Amendments to the Specification:

Please replace the paragraph beginning at page 6, line 15, with the following rewritten paragraph:

-- Particularly conserved regions and amino acid residues common to nectin polypeptides were identified by aligning nectin polypeptide sequences with each other and additional closely-related members of the nectin-Ig superfamily of proteins. The amino acid sequence of nectin-3 $\alpha$  and nectin-4 (SEQ ID Nos: 6 and 24) were compared with the amino acid sequences of other nectin and Ig family members (SEQ ID NO:20, 22, and 25), using a multiple sequence alignment program. The alignment of these sequences is shown in Table 2, and includes consensus residues (capitalized), which are identical among at least a majority [(three)] of the [five] amino acid sequences in the alignment. [In addition, lower case residues are shown on a separate line of Table 2 and represent residues that are not consensus residues, but are identical between human nectin-3 $\alpha$  and human nectin-4 (SEQ ID Nos: 6 and 24).] --

Please replace Table 2 beginning at page 7, line 7, with the following rewritten Table:

--

Table 2  
Conserved Nectin Amino Acids

(Hs=Homo sapiens)  
(Mus=Murine)

HUNECTION2 (SEQ ID NO:22)	~~~~~	~~~~~	MARAAALLPS	RSPPTPLLWP	LLLLLLL...
HUCD155 (SEQ ID NO:25)	~~~~~	~~~~~	~~~~~	MARAMAAWP	LLLVALLVLS
HUNECTION1 (SEQ ID NO:20)	~~~~~	~~~~~	~~~~~MARMG	LAGAAGRWWG	L...ALGLTA
HUNECTION3 (SEQ ID NO:6)	MARTLRPSPL	CPGGGKAQLS	SASLLGAGLL	LQPPTPPPLL	LLLFPLLFS
HUNECTION4 (SEQ ID NO:24)	~~~~~	~~~~~	~~~~~MPLSLG	AEMWGPEAWL	LLLLLLASFT
consensus				P W	LLL LL
HUNECTION2	51				100
HUCD155	..ETGAQDVR	VQVLPEVRGQ	LGGTVELPCH	L.LPPVPGLY	ISLVTWQRPD
HUNECTION1	WPPPGTGDDV	VQAPTQVPGF	LGDSVTLPCY	LQVPNMEVTH	VSQLTWAR..
HUNECTION3	FFLPGVHSQV	VQVNDSMYGF	IGTDVVLHCS	FANP.LPSVK	ITQVTWQK.S
HUNECTION4	RLCGALAGP.	IIVEPHVTAV	WGKNVSLKCL	I..EV..NET	ITQISWEKIH
	GRCP..AGE.	LETSDVVTVV	LGQDAKLPCF	YRGDS..GEQ	VGQVAWARVD
	<del>ePG ag</del>	<del>VQV</del>	<del>VtGv</del>	<del>LG V LPC</del>	<del>P e</del>
	<del>PG</del>	<del>VOV</del>	<del>V G</del>	<del>LG V LPC</del>	<del>P</del>
					I QV W R
					I QVTW R
HUNECTION2	101				150
HUCD155	APANHQNVA	FHPKMGPSFP	SPKPGSERLS	FVSAKQSTGQ	DTEAELQDAT
HUNECTION1	.HGEGSMAV	FHQTOGQPSYS	E....SKRLE	FVAARLG...	...AELRNAS
HUNECTION3	TNGSKQNVAI	YNPSMGVSV.	.LAPYRERVE	FL.....	...RPSFTDGT
HUNECTION4	.GKSSQTVAV	HHPQYGFVSQ	..GEYQGRVL	FKNYSLN...	.....DAT
	AGEGAQELAL	LHISKYGLHVS	..PAYEGRVE	QPPPPRNPL.	.....DGS
	<del>g</del>	<del>Q A</del>	<del>H yG SV</del>	<del>Y gRVE F</del>	<del>n</del>
	<del>Q VA</del>	<del>HP G SV</del>	<del>Y RVE F</del>	<del>F</del>	<del>DAT</del>
					DAT

HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	151 LALHGLTVED EGNITCEFAT FPKGSVRGMT WLRVIAKPKN QAEAQKVTF. LRMFGLRVED EGNITCLFVT FPQGSRSVDI WLRVLAKPQN TAEVQKVQL. IRLSRLELED EGVYICEFAT FPTGNRESQL NLTVMAKPTN WIEGTQAVLR ITLHNIGFSD SGKYICKAVT FPLGNAQSST TVTVLVEPTV SLIKGPDSLI VLLRNAVQAD EGEYECRVST FPAGSFQARL RLRVLVPPLP SLNPGP.ALE <del>L nL ED EG Y C F T FP GS q</del> <del>LRVLAKP N s E</del> <del>L</del> <del>L L ED EG Y C F T FP GS</del> <del>LRVLAKP N E</del> <del>L</del>	200
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	201 ....SQDPTT VALCISKEGR PPARISWLSS LDWEAKETQV SGTLAGTVTV ....TGEPVP MARCVSTGGR PPAQITWHS D LGGMPNTSQV PGFLSGTVTV AKKGQDDKVL VATCTSANGK PPSVSVSWETR LKGEARVPGD SGTPMAPVTV DGGNE...TV AAICIAATGK PVAHIDWEGD LGEM..ESTT TSFPNETATI EGQGL...TL AASC.TAEGS PAPSVTWDE VKGT..TSSR SFKHSRSAAV <del>g</del> <del>T aa C Sa G PPA I W L G S</del> <del>SG TVTV</del> <del>T A C SA G PPA I W L G S</del> <del>SG TVTV</del>	250
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	251 TSRFTLVPSG RADGVTVTCK VEH..ESFEE PALIPVTL SV RYPPEVSISG TSLWILVPSS QVDGKNVTCK VEH..ESFEK PQLLTVNLT V YYPPEVSISG ISRYRLVPSR EAHQQSLACI VNYHMDRFKE ....SLTLNV QYEPEVTIEG ISQYKLPFTR FARGRRITCV VKHP..ALEK DIRYSFILDY QYAPESVTVG TSEFHLVPSR SMNGQPLTCV VSHP..GLLQ DQRITHILHV SFLAEASVRG <del>TS LVP SR A G TC V H P FE d r i l v y PEVSI G</del> <del>TS LVP SR A G TC V H FE L V Y PEVSI G</del>	300
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	301 Y.DDN.WYLG RTDATLSCDV RSNPEPTGYD WSTTSGTFPT SAVAQGSQLV Y.DNN.WYLG QNEATLTCD A RSNPEPTGYN WSTTMGPLPP FAVAQAQLL F.DGN.WYLG RMDVKLTCKA DANPPATEYH WTLNLSLPK GVEAQNRTLF Y.DGN.WFVG RKGVNLCNA DANPPPFKSV WSRLDGQWPD GLLASDNTLH LEDQNLWHIG REGAMLKCLS EGQPPPSYN. WTRLDGPLPS GVRVDGDTLG <del>Y D N WYLG R g A L k C A NPPPT Y WSTL G LP G AQG TL</del> <del>Y D N WYLG R A L C A NPPPT Y WSTL G LP G AQG TL</del>	350
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	351 IH.AVDSL FN TTFVCTVTNA VGMGRAEQVI FVRETP.... IR.PVDKPIN TTLICNV TNA LGARQAELTV QVKEGP.... FKGPINYS LA GTYICEATNP IGTRSGQVEV NITEFPYTPS FVHPLTFNYS GVIYICKVTNS LGQRSDQKVI YISDPPTTTT LQPTIQWHPS F.PPLTTEHS GIYVCHVSNE FSSRDSQVTV DVLDPQEDSG KQ..... <del>F P l t s G YIC VTN G R Q V E P P q</del> <del>F P GTYIC VTN G R Q V Y E P</del>	400
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	401 .....RAS P...RDV..G PLVWGA VGGT LLVLLL LAGG .....PSE H...SGISRN AIIFLV LG.. ILVFLILLGI .....PPE HGRRAGPVPT AIIGGVAGSI LLVLIVVGGI TADIEDLATE PKKL PPLST LATIKDDTIA TIASVVGGA LFIVLVSVLA ....VDLV... ..SAS VVVVGVI AAL LFCLLVVVVV <del>d</del> <del>II GV G LLVLLV vG</del> <del>II GV G LLVLLV G</del>	450
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	451 SLAFILLVR RR....RKS .PGGAGGGAS GDGGFYDPKA QVLGN G D P V F GIYFYWSKCS REVLWHCHLC .PSSEHHQSC RN~~~~~ VVALRRRRHT FKGDYSTKKH .VYNGYSKA GIPQHPPMA QNLQY P D S D GIFCYRRRRT FRGDYFAKNY IPPSDMQKES QIDVLQOQDEL D..SYP.DSV LMSRYHRR... .KAQOMTQKY EEELTLTREN SIRRLHSHHT DPRSQPEESV <del>y RR y P e I h d L s P D s v</del> <del>Y RR P I I P S</del>	500

HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	501 WTPVVPGPME P.DGKDEEEE EEEKAEEKGL MLPPPPALED DMESQLDGSL ~~~~~ .DEKKAGPLG G.SSYEEEEEE EEEGGGGGER KVGPHPKYD EDAKRPYFTV .KKENKNP.. .VNNLIRKDY LEEPEKTQWN NVENLNRFER PMDYEDLKM GLRAEGHPDS LKDNSSCSVM SEEPEGRSYS TLTTVREIET QT...ELLSP ----- P n EePe e l P EE E
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	551 ISRRVYV~ ~~~~~ ~~~~~ ~~~~~ ~~~~~ ~~~~~ DEAEARQDGY GDRTLGYQYD PEQLDLAENM VSQNDGSFIS KKEWYV~ GM.KFVSDEH YDENEDDLVS HV...DGSVI SR...REWYV ~~~~~ GSGRAEEEEED QDEGIKQAMN HFVQENGLR AKPTGNGIYI NGRGHLV g A E DE I H g y A D

--